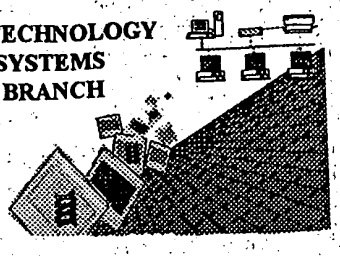
647 ALE

BIOTECHNOLOGY SYSTEMS BRANCH



RAW SEQUENCE LISTING ERROR REPORT

#9

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/430,412
Source: 1600
Date Processed by STIC: 11/20/2002

RECEIVED
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TECH CENTER 1600/2900

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
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Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, **EFS Submission User Manual** - ePAVE)
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3. **Hand Carry directly to:**
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Revised 01/29/2002

1600

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/430,412

DATE: 11/20/2002
TIME: 16:04:39

Input Set : A:\amended.txt
Output Set: N:\CRF4\11202002\I430412.raw

3 <110> APPLICANT: University of Ottawa
5 <120> TITLE OF INVENTION: Mutations of the 5' region of the human 5-HT1A gene,
6 associated proteins of the 5' region and a diagnostic
7 test for major depression and related mental illnesses
9 <130> FILE REFERENCE: 881014US1
11 <140> CURRENT APPLICATION NUMBER: 09/430,412
12 <141> CURRENT FILING DATE: 1999-10-29
14 <150> PRIOR APPLICATION NUMBER: 60/106,375
15 <151> PRIOR FILING DATE: 1998-10-30
17 <160> NUMBER OF SEQ ID NOS: 7
19 <170> SOFTWARE: PatentIn Ver. 2.1

Does Not Comply
Corrected Diskette Needed

ERRORED SEQUENCES

80 <210> SEQ ID NO: 2
81 <211> LENGTH: 24
82 <212> TYPE: DNA
83 <213> ORGANISM: human
85 <400> SEQUENCE: 2

E--> 86 aacgaagach nnnnnngtct tctt 24

see p.2 for error explanation

VARIABLE LOCATION SUMMARY
PATENT APPLICATION: US/09/430,412

DATE: 11/20/2002
TIME: 16:04:40

Input Set : A:\amended.txt
Output Set: N:\CRF4\11202002\I430412.raw

Use of n's or Xaa's(NEW RULES):
Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#:2; N Pos. 10,11,12,13,14,15,16

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/430,412

DATE: 11/20/2002
TIME: 16:04:40

Input Set : A:\amended.txt
Output Set: N:\CRF4\11202002\I430412.raw

L:86 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:2